PCT09

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         RAW SEQUENCE LISTING                DATE: 02/08/2002
         PATENT APPLICATION:  US/09/889,746  TIME: 11:18:39

         Input Set : A:\seqlist.txt
         Output Set: N:\CRF3\02082002\I889746.raw

      4 <110> APPLICANT: Ruelle, Jean-Louis
      6 <120> TITLE OF INVENTION: Novel Compounds
      9 <130> FILE REFERENCE: BM45352
     11 <140> CURRENT APPLICATION NUMBER: 09/889,746
C--> 12 <141> CURRENT FILING DATE: 2001-12-31
     14 <150> PRIOR APPLICATION NUMBER: 9901368.2
     15 <151> PRIOR FILING DATE: 1999-01-22
     17 <150> PRIOR APPLICATION NUMBER: 9901944.0
     18 <151> PRIOR FILING DATE: 1999-01-28
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     23 <150> PRIOR APPLICATION NUMBER: 9903417.5
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     26 <150> PRIOR APPLICATION NUMBER: 9903535.4
     27 <151> PRIOR FILING DATE: 1999-02-16
     29 <150> PRIOR APPLICATION NUMBER: PCT/EP00/00428
     30 <151> PRIOR FILING DATE: 2000-01-19
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     38 <212> TYPE: DNA
     39 <213> ORGANISM: Neisseria meningitidis
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     44 attaccattg ataagagcgg tatggcactt gccaatcgta tcacgcaaat gccccatacc   180
     45 accaaagtta tttatgaaga gcaaattcaa gagcaagcaa caggctctcg acagcttgcc   240
     46 gatgtgatgg cacagctcat tccaagtttg ggggtaagta gtggcactac cagtaacttt   300
     47 gggcaaacca tgcacggtcg tcaagtgcaa tttttgttaa atggcgtgcc tttgacaggt   360
     48 tcgcgagaca tctctagaca gcttaatagt atcaatccca atcaagtggc tagaattgaa   420
     49 gttttatcag gagcaaccag tatttatggg tctggagcaa caggcggttt gattaatatc   480
     50 gttactaagt ctgatttgga agaggagcaa tttgaaaccc gcatcggtgt acatggtagt   540
     51 aaattatcca gtgaaggtat cggttatcag gtaggtcaga gtgtagcagg tgtcagcgaa   600
     52 aatggtaatg tccttgcacg acttgatgtc gactatcgca ccacaggagg ggcatttgat   660
     53 gctaacggta aacgcatcgc tcctgagcct gcccaaactg ataagcaaga cagcaaaagc   720
     54 ctaagtgtca atacaaatgt tgattggcaa cttgacgaca agcaaaatat caatctggca   780
     55 ttgacgcatt ataacgacaa acaagatacc gattatgcac ctgattatgg taatcgcctt   840
     56 gcggtgttgt ttggagaaaa gccttcatta aatgccatca aaggcttatc attatcagaa   900
     57 cagccaaaaa ccaccaaaag cacctttaat atcaactatc atcatgatga tttgtggggt   960
     58 aacaccatca ataccaatgc ttattatcgc agagagaaag gcagatttta tcccttttgtt  1020
     59 gccccgtttt cgatcgccaa agccctgcct attttacaaa gcatgaattt gccatcagcc  1080
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66 <212> TYPE: PRT
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73                 20                  25                  30
74 Thr Val Glu Leu Glu Pro Val Val Ile Thr Ile Asp Lys Ser Gly Met
75             35                  40                  45
76 Ala Leu Ala Asn Arg Ile Thr Gln Met Pro His Thr Thr Lys Val Ile
77     50                  55                  60
78 Tyr Glu Glu Gln Ile Gln Glu Gln Ala Thr Gly Ser Arg Gln Leu Ala
79 65                  70                  75                  80
80 Asp Val Met Ala Gln Leu Ile Pro Ser Leu Gly Val Ser Ser Gly Thr
81                 85                  90                  95
82 Thr Ser Asn Phe Gly Gln Thr Met His Gly Arg Gln Val Gln Phe Leu
83                 100                 105                 110
84 Leu Asn Gly Val Pro Leu Thr Gly Ser Arg Asp Ile Ser Arg Gln Leu
85             115                 120                 125
86 Asn Ser Ile Asn Pro Asn Gln Val Ala Arg Ile Glu Val Leu Ser Gly
87     130                 135                 140
88 Ala Thr Ser Ile Tyr Gly Ser Gly Ala Thr Gly Gly Leu Ile Asn Ile
89 145                 150                 155                 160
90 Val Thr Lys Ser Asp Leu Glu Glu Glu Gln Phe Glu Thr Arg Ile Gly
91             165                 170                 175
92 Val His Gly Ser Lys Leu Ser Ser Glu Gly Ile Gly Tyr Gln Val Gly
93             180                 185                 190
94 Gln Ser Val Ala Gly Val Ser Glu Asn Gly Asn Val Leu Ala Arg Leu
95     195                 200                 205
96 Asp Val Asp Tyr Arg Thr Thr Gly Gly Ala Phe Asp Ala Asn Gly Lys
97     210                 215                 220
98 Arg Ile Ala Pro Glu Pro Ala Gln Thr Asp Lys Gln Asp Ser Lys Ser
99 225                 230                 235                 240
100 Leu Ser Val Asn Thr Asn Val Asp Trp Gln Leu Asp Asp Lys Gln Asn
101                 245                 250                 255
102 Ile Asn Leu Ala Leu Thr His Tyr Asn Asp Lys Gln Asp Thr Asp Tyr
103                 260                 265                 270
104 Ala Pro Asp Tyr Gly Asn Arg Leu Ala Val Leu Phe Gly Glu Lys Pro
105             275                 280                 285
106 Ser Leu Asn Ala Ile Lys Gly Leu Ser Leu Ser Glu Gln Pro Lys Thr
107     290                 295                 300
108 Thr Lys Ser Thr Phe Asn Ile Asn Tyr His His Asp Asp Leu Trp Gly
109 305                 310                 315                 320
110 Asn Thr Ile Asn Thr Asn Ala Tyr Tyr Arg Arg Glu Lys Gly Arg Phe
111                 325                 330                 335
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112 Tyr Pro Phe Val Ala Pro Phe Ser Ile Ala Lys Ala Leu Pro Ile Leu
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114 Gln Ser Met Asn Leu Pro Ser Ala Thr Leu Asp Ala Tyr Thr Lys Ala
115         355             360             365
116 Pro Gln Ala Arg Ala Tyr Gly Val Leu Gln Ser Glu Ser Lys Ala Glu
117     370             375             380
118 Val Leu Gly Arg Val Pro Asn Leu Asn Lys Pro Lys Arg Ala Leu Phe
119 385             390             395             400
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122 <211> LENGTH: 2409
123 <212> TYPE: DNA
124 <213> ORGANISM: Neisseria meningitidis
126 <400> SEQUENCE: 3
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128 cattgcgccg ccgccgatgc cgttgcggcg gaggaaacgg acaatccgac cgccggagga     120
129 agtgttcgga gcgtgtccga acccatgcag cctgccggcc tgagcctcgg ttcgacctgc     180
130 ctgttttgca gtaacgaaag cggcaaaccc gaaaaaaccg aatctgccgt caaaggaagc     240
131 ggcgaagggc ctgtgcccga aaaccacacg cgaattgtcg ccgacaaggt ggaagggcag     300
132 tcgcaggtca aggtacgcgc ggagggcggc gtcgttgtcg aacgcaaccg gacgaccctt     360
133 aatgccgact gggcggatta cgaccagtcg ggcgacaccg ttaccgtagg cgaccggttc     420
134 gccctccaac aggacggtac gctgattcgg ggcgaaaccc tgacctacaa tctcgagcag     480
135 cagaccggcg aagcgcacaa cgtccgcatg gaaaccgaac aaggcggacg gcggctgcaa     540
136 agcgtcagcc gcaccgccga aatgttgggc gaagggcatt acaaactgac ggaaacccaa     600
137 ttcaacacct gttccgccgg cgatgccggc tggtatgtca aggcagcctc tgtcgaagcc     660
138 gatcgggaaa aaggcatagg cgttgccaaa cacgccgcct tcgtgttcgg cggcgttcct     720
139 attttctaca cccttgggc ggacttcccg cttgacggca accgcaaaag cggcctgctt     780
140 gttccctcac tgtccgccgg ttcggacggc gtttcccttt ccgttcccta ttatttcaac     840
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142 tttgacgggc aggtacgcta cctgcggccg gattatgccg gccagtccga cctgacctgg     960
143 ctgccgcacg acaagaaaag cggcaggaat aaccgctatc aggcgaaatg gcagcatcgg    1020
144 cacgacattt ccgacacgct tcaggcgggt gtcgatttca accaagtctc cgacagcggc    1080
145 tactaccgcg acttttacgg caacaaagaa atcgccggca acgtcaacct caaccgccgt    1140
146 gtatggctgg attatggcgg cagggcggcg ggcggcagcc tgaatgccgg cctttcggtt    1200
147 ctgaaatacc agacgctggc aaaccaaagc ggctacaaag acaaaccgta tgccctgatg    1260
148 ccgcgccttt ccgccgattg cgcaaaaac accggcaggg cgcaaatcgg cgtgtccgca    1320
149 caatttaccc gcttcagcca cgacagccgc caagacggca gccgcctcgt cgtctatccc    1380
150 gacatcaaat gggatttcag caacagctgg ggttacgtcc gtcccaaact cggactgcac    1440
151 gccacctatt acagcctcaa ccgcttcggc agccaagaag cccgacgcgt cagccgcact    1500
152 ctacccatcg tcaacatcga cagcggcatg accttcgaac gcaatacgcg gatgttcggc    1560
153 ggagaagtcc tgcaaaccct cgagccgcgc ctgttctaca actatattcc tgccaaatcc    1620
154 caaaacgacc tgcccaattt tgattcgtcg gaaagcagct cggctacgg gcagctttt    1680
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156 gtgcaaagcc gtattttgga cggcgcgacg ggggcagagc gtttccgcgc cggcatcggg    1800
157 cagaaattct acttcaaaaa cgacgcagtc atgcttgacg gcagtgtcgg caaaaaaccg    1860
158 cgcagccgtt ccgactgggt ggcattcgcc tccagcggca tcggcagccg cttcatcctc    1920
159 gacagcagca tccactacaa ccaaaacgac aaacgcgccg agaactacgc cgtcggtgca    1980
160 agctaccgtc ccgcacaggg caaagtgctg aacgcccgct acaaatacgg cgcaacgaa    2040
161 aaaatctacc tgaagtccga cggttcctat ttttacgaca aactcagcca gctcgacctg    2100
162 tccgcacaat ggcccctgac gcgcaacctg tcggccgtcg tccgttacaa ctacggtttt    2160
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165 gtcttttct cacttcagtt gaaagacctc agcagtgtcg gcagaaaccc cgcagacagg    2340
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170 <211> LENGTH: 802
171 <212> TYPE: PRT
172 <213> ORGANISM: Neisseria meningitidis
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178             20                  25                  30
179 Thr Asp Asn Pro Thr Ala Gly Gly Ser Val Arg Ser Val Ser Glu Pro
180         35                  40                  45
181 Met Gln Pro Ala Gly Leu Ser Leu Gly Ser Thr Cys Leu Phe Cys Ser
182     50                  55                  60
183 Asn Glu Ser Gly Lys Pro Glu Lys Thr Glu Ser Ala Val Lys Gly Ser
184 65                  70                  75                  80
185 Gly Glu Gly Pro Val Pro Glu Asn His Thr Arg Ile Val Ala Asp Lys
186                 85                  90                  95
187 Val Glu Gly Gln Ser Gln Val Lys Val Arg Ala Glu Gly Gly Val Val
188                 100                 105                 110
189 Val Glu Arg Asn Arg Thr Thr Leu Asn Ala Asp Trp Ala Asp Tyr Asp
190         115                 120                 125
191 Gln Ser Gly Asp Thr Val Thr Val Gly Asp Arg Phe Ala Leu Gln Gln
192     130                 135                 140
193 Asp Gly Thr Leu Ile Arg Gly Glu Thr Leu Thr Tyr Asn Leu Glu Gln
194 145                 150                 155                 160
195 Gln Thr Gly Glu Ala His Asn Val Arg Met Glu Thr Glu Gln Gly Gly
196                 165                 170                 175
197 Arg Arg Leu Gln Ser Val Ser Arg Thr Ala Glu Met Leu Gly Glu Gly
198             180                 185                 190
199 His Tyr Lys Leu Thr Glu Thr Gln Phe Asn Thr Cys Ser Ala Gly Asp
200             195                 200                 205
201 Ala Gly Trp Tyr Val Lys Ala Ala Ser Val Glu Ala Asp Arg Glu Lys
202     210                 215                 220
203 Gly Ile Gly Val Ala Lys His Ala Ala Phe Val Phe Gly Gly Val Pro
204 225                 230                 235                 240
205 Ile Phe Tyr Thr Pro Trp Ala Asp Phe Pro Leu Asp Gly Asn Arg Lys
206                 245                 250                 255
207 Ser Gly Leu Leu Val Pro Ser Leu Ser Ala Gly Ser Asp Gly Val Ser
208             260                 265                 270
209 Leu Ser Val Pro Tyr Tyr Phe Asn Leu Ala Pro Asn Leu Asp Ala Thr
210         275                 280                 285
211 Phe Ala Pro Ser Val Ile Gly Glu Arg Gly Ala Val Phe Asp Gly Gln
212     290                 295                 300
213 Val Arg Tyr Leu Arg Pro Asp Tyr Ala Gly Gln Ser Asp Leu Thr Trp
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RAW SEQUENCE LISTING   DATE: 02/08/2002
PATENT APPLICATION: US/09/889,746   TIME: 11:18:39

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\02082002\I889746.raw

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214 305                    310                    315                     320
215 Leu Pro His Asp Lys Lys Ser Gly Arg Asn Asn Arg Tyr Gln Ala Lys
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217 Trp Gln His Arg His Asp Ile Ser Asp Thr Leu Gln Ala Gly Val Asp
218             340                    345                     350
219 Phe Asn Gln Val Ser Asp Ser Gly Tyr Tyr Arg Asp Phe Tyr Gly Asn
220             355                    360                     365
221 Lys Glu Ile Ala Gly Asn Val Asn Leu Asn Arg Arg Val Trp Leu Asp
222       370                    375                    380
223 Tyr Gly Gly Arg Ala Ala Gly Gly Ser Leu Asn Ala Gly Leu Ser Val
224 385                    390                    395                     400
225 Leu Lys Tyr Gln Thr Leu Ala Asn Gln Ser Gly Tyr Lys Asp Lys Pro
226             405                    410                     415
227 Tyr Ala Leu Met Pro Arg Leu Ser Ala Asp Trp Arg Lys Asn Thr Gly
228             420                    425                     430
229 Arg Ala Gln Ile Gly Val Ser Ala Gln Phe Thr Arg Phe Ser His Asp
230             435                    440                     445
231 Ser Arg Gln Asp Gly Ser Arg Leu Val Val Tyr Pro Asp Ile Lys Trp
232       450                    455                    460
233 Asp Phe Ser Asn Ser Trp Gly Tyr Val Arg Pro Lys Leu Gly Leu His
234 465                    470                    475                     480
235 Ala Thr Tyr Tyr Ser Leu Asn Arg Phe Gly Ser Gln Glu Ala Arg Arg
236             485                    490                     495
237 Val Ser Arg Thr Leu Pro Ile Val Asn Ile Asp Ser Gly Met Thr Phe
238             500                    505                     510
239 Glu Arg Asn Thr Arg Met Phe Gly Gly Glu Val Leu Gln Thr Leu Glu
240       515                    520                    525
241 Pro Arg Leu Phe Tyr Asn Tyr Ile Pro Ala Lys Ser Gln Asn Asp Leu
242       530                    535                    540
243 Pro Asn Phe Asp Ser Ser Glu Ser Ser Phe Gly Tyr Gly Gln Leu Phe
244 545                    550                    555                     560
245 Arg Glu Asn Leu Tyr Tyr Gly Asn Asp Arg Ile Asn Thr Ala Asn Ser
246             565                    570                     575
247 Leu Ser Ala Ala Val Gln Ser Arg Ile Leu Asp Gly Ala Thr Gly Ala
248             580                    585                     590
249 Glu Arg Phe Arg Ala Gly Ile Gly Gln Lys Phe Tyr Phe Lys Asn Asp
250       595                    600                    605
251 Ala Val Met Leu Asp Gly Ser Val Gly Lys Lys Pro Arg Ser Arg Ser
252       610                    615                    620
253 Asp Trp Val Ala Phe Ala Ser Ser Gly Ile Gly Ser Arg Phe Ile Leu
254 625                    630                    635                     640
255 Asp Ser Ser Ile His Tyr Asn Gln Asn Asp Lys Arg Ala Glu Asn Tyr
256             645                    650                     655
257 Ala Val Gly Ala Ser Tyr Arg Pro Ala Gln Gly Lys Val Leu Asn Ala
258             660                    665                     670
259 Arg Tyr Lys Tyr Gly Arg Asn Glu Lys Ile Tyr Leu Lys Ser Asp Gly
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261 Ser Tyr Phe Tyr Asp Lys Leu Ser Gln Leu Asp Leu Ser Ala Gln Trp
262       690                    695                    700
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VERIFICATION SUMMARY   DATE: 02/08/2002
PATENT APPLICATION: US/09/889,746   TIME: 11:18:40

Input Set : **A:\seqlist.txt**
Output Set: **N:\CRF3\02082002\I889746.raw**

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date